



STIC Search Report

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TO: Karen A Lacourciere
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Art Unit: 1635
Thursday, July 22, 2004

Case Serial Number: 10/006366

From: David Schreiber
Location: Biotech-Chem Library
Remsen E01A61
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Search Notes

S8 31 RD (unique items)
?show files;ds;t/3,k/all
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 (c) 1998 Inst for Sci Info
File 40:Enviroline(R) 1975-2004/Jul
File 50:CAB Abstracts 1972-2004/Aug
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File 103:Energy SciTec 1974-2004/Aug B2
 (c) 2004 Contains copyrighted material
File 156>ToxFile 1965-2004/Sep W2
 (c) format only 2004 The Dialog Corporation
File 162:Global Health 1983-2004/Aug
 (c) 2004 CAB International
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 (c) 2004 Royal Soc Chemistry
File 35:Dissertation Abs Online 1861-2004/Aug
 (c) 2004 ProQuest Info&Learning
File 48:SPORTDiscus 1962-2004/Sep
 (c) 2004 Sport Information Resource Centre
File 91:MANTIS(TM) 1880-2004/Sep
 2001 (c) Action Potential
File 149:TGG Health&Wellness DB(SM) 1976-2004/Aug W4
 (c) 2004 The Gale Group

File 159:Cancerlit 1975-2002/Oct
(c) format only 2002 Dialog Corporation
File 164:Allied & Complementary Medicine 1984-2004/Sep
(c) 2004 BLHCIS
File 444:New England Journal of Med. 1985-2004/Sep W2
(c) 2004 Mass. Med. Soc.
File 467:ExtraMED(tm) 2000/Dec
(c) 2001 Informania Ltd.

Set	Items	Description
S1	0	(CLASS II ACTIVATOR) (S) (ANTISENSE OR RIBOZYME?)
S2	0	(CLASS II ACTIVATOR) AND (ANTISENSE OR RIBOZYME?)
S3	2755	CIITA
S4	31	S3 (S) (ANTISENSE OR RIBOZYME?)
S5	21	RD (unique items)
S6	2756	(CLASS II ACTIVATOR?) OR (CIITA)
S7	62	S6 AND (VARIANT? OR MHC2TA OR MHC2TAII OR MHC2TAIII OR MGC- 2TAIV OR MHC2TAV OR MHC2TAVI)
S8	31	RD (unique items)

RESULT 6		1	(bases 1 to 43)
LOCUS	AX523241	49 bp	DNA
DEFINITION	Sequence 911 from Patent WO20064731.	linear	PAT 24-OCT-2002
VERSION	AX523241		
JOURNAL	TELMERMAN, A., Amson, R., Tuijnder, M. and Sussini, L.		
FEATURES	Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as medicines		
SOURCE	Homo sapiens (human)		
ORGANISM	Homo sapiens		
REFERENCE	Ting, J.-Pan, Yung, and Chin, K.-C.		
AUTHORS	Forms of class II MHC transactivator (CLITA)		
TITLE	Patent: US 5994505-A 7 30-NOV-1999;		
JOURNAL/Qualifiers	Location/Qualifiers 1..43		
FEATURES	Source /organism="unknown" /mol_type="unassigned DNA"		
SOURCE			
RESULT 7		1	(bases 1 to 43)
LOCUS	AR290618	47 bp	DNA
DEFINITION	Sequence 2353 from patent US 6537751.	linear	PAT 12-JUN-2003
VERSION	AR290618		
JOURNAL	Cohen,D., Chumakov,I. and Blumenfeld,M.		
FEATURES	Biallelic markers for use in constructing a high density disequilibrium map of the human genome		
SOURCE	Patent: US 6537751-A 23-3 25-MAR-2003; Location/Qualifiers 1..47		
ORGANISM	Unknown.		
REFERENCE	Chumakov,I. and Blumenfeld,M.		
AUTHORS	Biallelic markers for use in constructing a high density disequilibrium map of the human genome		
TITLE	Patent: US 6537751-A 23-3 25-MAR-2003;		
JOURNAL/Qualifiers	Location/Qualifiers 1..47		
FEATURES	/organism="unknown" /mol_type="genomic DNA"		
SOURCE			
RESULT 8		1	(bases 1 to 43)
LOCUS	AR091697	43 bp	DNA
DEFINITION	Sequence 7 from patent US 5994505.	linear	PAT 07-SEP-2000
VERSION	AR091697		
JOURNAL	Unknown.		
FEATURES	Unknown.		
SOURCE	Unknown.		
RESULT 9		1	(bases 1 to 43)
LOCUS	AR091698/c	43 bp	DNA
DEFINITION	Sequence 8 from patent US 5994505.	linear	PAT 07-SEP-2000
VERSION	AR091698		
JOURNAL	Unknown.		
FEATURES	Unknown.		
SOURCE	Unknown.		
RESULT 10		1	(bases 1 to 43)
LOCUS	AR290806/c	47 bp	DNA
DEFINITION	Sequence 2541 from patent US 6537751.	linear	PAT 12-JUN-2003
VERSION	AR290806		
JOURNAL	Cohen,D., Chumakov,I. and Blumenfeld,M.		
FEATURES	Biallelic markers for use in constructing a high density disequilibrium map of the human genome		
SOURCE	Patent: US 6537751-A 25-4 25-MAR-2003; Location/Qualifiers 1..47		
ORGANISM	Unknown.		
REFERENCE	Chumakov,I. and Blumenfeld,M.		
AUTHORS	Biallelic markers for use in constructing a high density disequilibrium map of the human genome		
TITLE	Patent: US 6537751-A 25-4 25-MAR-2003;		
JOURNAL/Qualifiers	Location/Qualifiers 1..47		
FEATURES	/organism="unknown" /mol_type="genomic DNA"		
SOURCE			
RESULT 11		1	(bases 1 to 43)
LOCUS	4166	4208	DNA
DEFINITION	CCCCAAGCTGCTGGATTCACAGCTGGCCATCTGCACGGGCC	linear	PAT 07-SEP-2000
VERSION	4166		
JOURNAL	Unknown.		
FEATURES	Unknown.		
SOURCE	Unknown.		
RESULT 12		1	(bases 1 to 43)
LOCUS	4167	4209	DNA
DEFINITION	CCCCAACCTCCCTGGATTACAGGCGTGTGCACTGGCCCACTGGCATCCGGCC	linear	PAT 07-SEP-2000
VERSION	4167		
JOURNAL	Unknown.		
FEATURES	Unknown.		
SOURCE	Unknown.		

TELEPHONE: 919-420-2200
 TELEFAX: 919-881-3175
 INFORMATION FOR SEQ ID NO: 2:
 LENGTH: 21 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 MOLECULE TYPE: linear
 DESCRIPTION: /desc = "DNA oligonucleotide"
 US-08-816-617A-2

RESULT 77
 US-08-816-617A-3/c
 Query Match 0.3%; Score 21; DB 1; Length 21;
 Best Local Similarity 100.0%; Pred. No. 80;
 Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 114 GCTGCGCTGGCTGGATCCCTA 134
 Db 1 GCTGCGCTGGCTGGATCCCTA 21

RESULT 77
 US-09-540-257B-1
 Query Match 0.3%; Score 21; DB 1; Length 21;
 Best Local Similarity 100.0%; Pred. No. 80;
 Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 373 AGCAGCGCTGGTGTGAGATG 393
 Db 1 AGCAGCGCTGGTGTGAGATG 21

RESULT 79
 US-09-018-637-27/C
 Sequence 27, Application US/09038637
 Patent No. 6235410
 GENERAL INFORMATION:
 APPLICANT: Ting, Jenny P.-Y.
 APPLICANT: Piskurich, Janet
 TITLE OF INVENTION: No. 6022741el Regulatory Genetic DNA that
 TITLE OF INVENTION: Regulates the Class II Transactivator (CIITA)
 NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:
 ADDRESSEE: Bell, Seitzer, Park & Gibson
 STREET: 1211 East Morehead Street
 CITY: Charlotte
 STATE: North Carolina
 COUNTRY: United States
 ZIP: 28234

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/816.617A
 FILING DATE:
 CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:
 NAME: Sibley, Kenneth D.
 REGISTRATION NUMBER: 31,665
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 919-420-2200
 TELEFAX: 919-881-3175
 SEQUENCE CHARACTERISTICS:
 LENGTH: 21 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 MOLECULE TYPE: linear
 DESCRIPTION: /desc = "DNA oligonucleotide"
 US-08-816-617A-3

Query Match 0.3%; Score 21; DB 1; Length 21;
 Best Local Similarity 100.0%; Pred. No. 80;
 Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 78
 US-09-540-257B-1
 Sequence 1, Application US/09540257B
 Patent No. 6518012
 GENERAL INFORMATION:
 APPLICANT: Tomasi, Thomas
 TITLE OF INVENTION: Method for Regulating the Expression of MHC Antigens and
 FILE REFERENCE: 03551.0048
 CURRENT APPLICATION NUMBER: US/09/540,257B
 CURRENT FILING DATE: 2000-03-31
 PRIOR APPLICATION NUMBER: US 60/146,275; US 60/127,591
 PRIOR FILING DATE: 1999-07-29; 1999-04-02
 NUMBER OF SEQ ID NOS: 26

SEQ ID NO: 1
 LENGTH: 21
 TYPE: DNA
 ORGANISM: Artificial Sequence
 FEATURE:
 OTHER INFORMATION: Forward primer for Human CIITA
 US-09-540-257B-1

Query Match 0.3%; Score 21; DB 1; Length 21;
 Best Local Similarity 100.0%; Pred. No. 80;
 Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 373 AGCAGCGCTGGTGTGAGATG 393
 Db 1 AGCAGCGCTGGTGTGAGATG 21

RESULT 79
 US-09-018-637-27/C
 Sequence 27, Application US/09038637
 Patent No. 6235410
 GENERAL INFORMATION:
 APPLICANT: Sidransky, David
 TITLE OF INVENTION: DETECTION OF NEOPLASM BY ANALYSIS OF SALIVA
 NUMBER OF SEQUENCES: 195
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Fish & Richardson P.C.
 STREET: 4225 Executive Square, Suite 1400
 CITY: La Jolla
 STATE: CA
 COUNTRY: USA
 ZIP: 92037

COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette
 COMPUTER: IBM Compatible
 OPERATING SYSTEM: Windows 95
 SOFTWARE: FastSEQ for Windows Version 2.0b
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/038,637
 FILING DATE: 10-MAR-1998
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/579,233
 FILING DATE: 28-DEC-1995
 APPLICATION NUMBER: 08/152,313
 FILING DATE: 12-NOV-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: Haile, Liba A.
 REGISTRATION NUMBER: 38,347
 REFERENCE/DOCKET NUMBER: 07265/146001
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 619/678-5070
 TELEFAX: 619/678-5099
 INFORMATION FOR SEQ ID NO: 27:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 24 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: Genomic DNA